The structures were solved by direct methods and refined by full-matrix least squares for all reflections. H atoms were placed geometrically (except those of the H₂O molecules in BRL-53888A, which were obtained from a difference Fourier synthesis) and refined with a riding model and with U_{iso} constrained to be $1.25U_{eq}$ of the parent atom.

For both compounds, data collection: XSCANS (Siemens, 1996); cell refinement: XSCANS; data reduction: XSCANS; program(s) used to solve structures: SIR92 (Altomare *et al.*, 1994); program(s) used to refine structures: SHELXL93 (Sheldrick, 1993); molecular graphics: DIAMOND (Bergerhoff, 1996); software used to prepare material for publication: PARST (Nardelli, 1983).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: BM1280). Services for accessing these data are described at the back of the journal.

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4-Ethyl-2,3-dihydro-4*H*-pyrido[3,2-*e*]-1,2,4thiadiazine 1,1-dioxide and 4-ethyl-2,3-dihydro-4*H*-pyrido[4,3-*e*]-1,2,4-thiadiazine 1,1-dioxide[†]

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Abstract

A series of 4H-1,2,4-pyridothiadiazine 1,1-dioxides and 2,3-dihydro-4H-1,2,4-pyridothiadiazine 1,1-dioxides were tested as possible allosteric modulators of the (R/S)-2-amino-3-(3-hydroxy-5-methylisoxazol-4-yl) propionic acid receptors; the most active is 4-ethyl-2,3dihydro-4H-pyrido[3,2-e]-1,2,4-thiadiazine 1,1-dioxide, $C_8H_{11}N_3O_2S$. Its crystal molecular geometry is compared with that of the -pyrido[4,3-e]- compound, $C_8H_{11}N_3O_2S$, a less potent analogue.

Comment

A series of 4H- and 2,3-dihydro-4H-1,2,4-pyridothiadiazine 1,1-dioxides, belonging to three different chemical classes (as a function of the N-atom position in the heterocycle) and bearing various alkyl and aryl substituents at the 2, 3 and 4 positions, were synthesized and tested as possible allosteric modulators of the (R/S)-2-amino-3-(3-hydroxy-5-methylisoxazol-4yl) propionic acid (AMPA) receptors. Many compounds were found to be more potent than the reference compounds diazoxide (Bandoli & Nicolini, 1977) and aniracetam as potentiators of the AMPA current in rat cortex mRNA-injected Xenopus oocytes. The most active compound, 4-ethyl-2,3-dihydro-4H-pyrido[3,2-e]-1,2,4-thiadiazine 1,1-dioxide, (1), revealed an in vitro activity not far from that of cyclothiazide, the most potent allosteric modulator of AMPA receptors reported to date. Structure-activity relationships were deduced and indicated the possible dissociation between the structure requirements leading to a biological activity

[†] Systematic names: 4-ethyl-3,4-dihydro-2*H*-pyrido[3,2-*e*]-1,2,4-thiadiazine 1,1-dioxide and 4-ethyl-3,4-dihydro-2*H*-pyrido[4,3-*e*]-1,2,4thiadiazine 1,1-dioxide.



of the pyridothiadiazines even on AMPA receptors or on ATP-sensitive K^+ channels (Pirotte *et al.*, 1998).

The crystal structure of (1) exhibits two independent conformers, (1A) and (1B), in the asymmetric unit which are distinguishable by the different orientations of the terminal methyl group of the side chains. N2 was found to be clearly of sp^3 pyramidal geometry [sums

of bond angles around N2 are 336 (2) and 328 (2)° for conformers (1A) and (1B), respectively] whereas N4 was found to be of sp^2 geometry [sums of bond angles around N4 are 359.3 (3) and 356.7 (3)° for (1A) and (1B), respectively].

4-Ethyl-2,3-dihydro-4*H*-pyrido[4,3-*e*]-1,2,4-thiadiazine 1,1-dioxide, (2), crystallizes with one molecule in the asymmetric unit geometrically superimposable to conformer (1*A*). N2 approximates the sp^3 pyramidal geometry [sum of angles is 337 (2)°] and N4 coincides with an sp^2 geometry [sum of angles is 360.0 (2)°]. Compound (2) shows a strong analogy with the previously reported 2,4,7-trimethyl-2,3-dihydro-4*H*-pyrido[4,3-*e*]-1,2,4-thiadiazinium 1,1-dioxide iodide (Dupont *et al.*,







Fig. 1. Molecular structure with atom-labelling scheme for (a) molecule (1A), (b) molecule (1B) and (c) molecule (2). Displacement ellipsoids are shown at 50% probability levels. H atoms are drawn as small circles of an arbitrary radius.

1995) bearing a methyl group on the N atom at the 2 position [angles around N2 and N4 are 341.8(9) and $359.6(9)^{\circ}$, respectively].

A comparison of the molecular geometries of (1) and (2) shows no significant variation. In particular, the sp^3 nature of N at the 2 position of the thiadiazine diazoxide ring is almost similar in both structures. Hence the difference in their biological activity is primarily due to the nitrogen position in the pyridine ring.

The shortest intermolecular contacts include NH on the 2 position in both structures (Tables 2 and 4). In (1A) and (2), the geometries agree with nearly 'linear' NH···O or NH···H hydrogen bonds. In (1B) the N12...O2 and N12...N9 intermolecular distances are close to the sum of their respective van der Waals radii. In this case, the occurrence of an asymmetric 'bifurcated' hydrogen bond may be tentatively suggested (Taylor et al., 1984).

Experimental

The compounds were synthesized at the Laboratory of Medicinal Chemistry of Liège (Pirotte et al., 1998). Crystals were obtained by slow evaporation of a methanol solution at room temperature.

Compound (1)

Crystal data

$C_8H_{11}N_3O_2S$	Cu $K\alpha$ radiation
$M_r = 213.26$	$\lambda = 1.54180 \text{ Å}$
Monoclinic	Cell parameters from 27
$P2_1/n$	reflections
a = 7.7385 (12) Å	$\theta = 26.14 - 37.34^{\circ}$
b = 12.199(3) Å	$\mu = 2.839 \text{ mm}^{-1}$
c = 20.667(2) Å	T = 293 (2) K
$\beta = 99.735(18)^{\circ}$	Prismatic
V = 1922.9 (6) Å ³	$0.46 \times 0.30 \times 0.23$ mm
Z = 8	Colourless
$D_x = 1.473 \text{ Mg m}^{-3}$	
D_m not measured	
Data collection	
Data contection	
Stoe-Siemens AED four-	1937 reflections with

circle diffractometer ω scans Absorption correction: semi-empirical, ψ scans (EMPIR; Stoe & Cie, 1987a) $T_{\rm min} = 0.294, T_{\rm max} = 0.341$ 3719 measured reflections 3451 independent reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.045$ $wR(F^2) = 0.132$

h $I > 2\sigma(I)$ $R_{\rm int} = 0.028$ $\theta_{\rm max} = 67.47^{\circ}$ $h = -9 \rightarrow 9$ $k=-3\rightarrow 14$ $l = -24 \rightarrow 24$

mm

2 standard reflections frequency: 60 min intensity decay: 4.1%

 $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.343 \ {\rm e} \ {\rm \AA}^{-3}$ $\Delta \rho_{\rm min} = -0.239 \ {\rm e} \ {\rm \AA}^{-3}$

S = 0.943	Extinction correction:
3451 reflections	SHELXL97 (Sheldrick,
262 parameters	1997a)
H atoms were restrained	Extinction coefficient:
(included as riding atoms)	0.0166 (8)
and atoms H2 and H12	Scattering factors from
were refined	International Tables for
$w = 1/[\sigma^2(F_o^2) + (0.0819P)^2]$	Crystallography (Vol. C)
where $P = (F_0^2 + 2F_c^2)/3$	

Table 1. Selected geometric parameters (Å, °) for (1)

	0	•	5
S1—N2	1.632 (3)	\$11—N12	1.627 (3)
N2—C3	1.432 (5)	N12-C13	1.462 (4)
N2—H2	0.87 (3)	N12—H12	0.88 (3)
C3—N4	1.443 (4)	C13—N14	1.452 (4)
N4—C5	1.376 (4)	N14-C15	1.372 (4)
N4—C11	1.466 (5)	N14C21	1.459 (4)
N2-S1-C10	102.06 (15)	N12-S11-C20	102.00 (14
C3-N2-S1	113.9 (2)	C13-N12-S11	112.9 (2)
C3—N2—H2	122 (2)	C13-N12-H12	113 (2)
S1-N2-H2	100 (2)	S11—N12—H12	103 (2)
N2-C3-N4	113.6 (3)	N14-C13-N12	111.9 (3)
C5—N4—C3	118.2 (3)	C15-N14-C13	117.8 (2)
C5-N4-C11	123.6 (3)	C15-N14-C21	121.5 (3)
C3-N4-C11	117.5 (3)	C13-N14-C21	117.4 (3)
N4-C11-C12	111.9 (3)	N14-C21-C22	114.0 (3)
C10-S1-N2-C3	42.7 (3)	C20-S11-N12-C13	-43.4 (2)
S1-N2-C3-N4	-65.5(4)	S11-N12-C13-N14	67.5 (3)
N2-C3-N4-C5	46.9 (4)	N12-C13-N14-C15	-52.7 (4)
N2-C3-N4-C11	-142.2(3)	N12-C13-N14-C21	147.4 (3)
C3-N4-C5-C10	-9.5(5)	C13-N14-C15-C20	17.6 (4)

Table 2. Hydrogen-bonding geometry $(Å, \circ)$ for (1)

-	0	00		
D — $H \cdots A$	D—H	H···A	$D \cdot \cdot \cdot A$	D — $\mathbf{H} \cdots A$
N2—H2···O12 ¹	0.87 (3)	2.25 (3)	3.034 (4)	149 (3)
$N12 - H12 \cdot \cdot \cdot O2^{ii}$	0.88 (3)	2.80 (3)	3.285 (3)	116 (3)
N12—H12···N9 ⁱⁱ	0.88 (3)	2.47 (3)	3.305 (4)	159 (3)
			,	

Symmetry codes: (i) 1 + x, y, z; (ii) $x - \frac{1}{2}, \frac{3}{2} - y, \frac{1}{2} + z$.

Compound (2)

Crysiai aala	
$C_8H_{11}N_3O_2S$	Cu $K\alpha$ radiation
$M_r = 213.26$	$\lambda = 1.54180 \text{ Å}$
Monoclinic	Cell parameters from 32
$P2_1/n$	reflections
a = 8.1628 (4) Å	$\theta = 29.96 - 37.23^{\circ}$
b = 11.6259 (8) Å	$\mu = 2.835 \text{ mm}^{-1}$
c = 10.2405 (11) Å	T = 293 (2) K
$\beta = 97.841 (7)^{\circ}$	Prismatic
$V = 962.74 (13) \text{ Å}^3$	$0.46 \times 0.42 \times 0.38$ mm
Z = 4	Colourless
$D_x = 1.471 \text{ Mg m}^{-3}$	
D_m not measured	

Data collection

Stoe-Siemens AED four-1387 reflections with $I > 2\sigma(I)$ circle diffractometer $R_{int} = 0.038$ ω scans $\theta_{\rm max} = 68.01^{\circ}$ Absorption correction: $h = -9 \rightarrow 9$ semi-empirical, ψ scans $k = -13 \rightarrow 0$ (EMPIR; Stoe & Cie, $l = 0 \rightarrow 12$ 1987a) 2 standard reflections $T_{\rm min} = 0.329, T_{\rm max} = 0.520$ 1803 measured reflections frequency: 60 min intensity decay: 2.6% 1707 independent reflections

Refinement

Refinement on F^2	$(\Delta/\sigma)_{\rm max} < 0.001$
$R[F^2 > 2\sigma(F^2)] = 0.038$	$\Delta \rho_{\rm max} = 0.215 \ {\rm e} \ {\rm \AA}^{-3}$
$wR(F^2) = 0.106$	$\Delta ho_{\min} = -0.265 \text{ e } \text{\AA}^{-3}$
S = 1.060	Extinction correction:
1707 reflections	SHELXL97 (Sheldrick,
132 parameters	1997a)
H atoms were restrained	Extinction coefficient:
(included as riding atoms)	0.0260 (14)
and atom H2 which was	Scattering factors from
refined	International Tables for
$w = 1/[\sigma^2(F_o^2) + (0.0469P)^2]$	Crystallography (Vol. C)
+ 0.5899 <i>P</i>]	
where $P = (F_o^2 + 2F_c^2)/3$	

Table 3. Selected geometric parameters (Å, $^{\circ}$) for (2)

\$1—N2	1.627 (2)	C3N4	1.463 (3)
N2C3	1.435 (3)	N4—C5	1.353 (3)
N2—H2	0.75 (3)	N4C11	1.465 (3)
N2-S1-C10	100.97 (11)	C5-N4-C3	121.4 (2)
C3-N2-S1	111.66 (17)	C5-N4-C11	122.9 (2)
C3—N2—H2	115 (2)	C3—N4—C11	115.7 (2)
\$1-N2-H2	110 (2)	N4-C11-C12	112.0 (2)
N2—C3—N4	113.7 (2)		
C10-S1-N2-C3	54.77 (19)	N2-C3-N4-C11	-150.9 (2)
S1-N2-C3-N4	-63.6(3)	C3-N4-C5-C10	1.7 (4)
N2-C3-N4-C5	32.5 (3)		

Table 4. Hydrogen-bonding geometry (Å, °) for (2)

For both compounds, data collection: *DIF*4 (Stoe & Cie, 1987b); cell refinement: *DIF*4; data reduction: *REDU*4 (Stoe & Cie, 1987c); program(s) used to solve structures: *SHELXS97* (Sheldrick, 1997b); program(s) used to refine structures: *SHELXL97* (Sheldrick, 1997a); molecular graphics: *ORTEPIII* (Burnett & Johnson, 1996); software used to prepare material for publication: *SHELXL97*.

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© 1999 International Union of Crystallography Printed in Great Britain – all rights reserved Sheldrick, G. M. (1997b). SHELXS97. Program for the Solution of Crystal Structures. University of Göttingen, Germany.

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2,3,7,8,12,13,17,18-Octafluoro-5,10,15,20tetrakis(pentafluorophenyl)porphyrin

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Abstract

The core of the title compound, $C_{44}H_2F_{28}N_4$, is essentially planar while the pentafluorophenyl groups are nearly perpendicular to the mean porphyrin plane. The molecule is centrosymmetric.

Comment

The title compound, F_{28} TPP, was prepared as part of a study of β -octafluoroporphyrins (Leroy *et al.*, 1997), a new class of highly electron-deficient ligands. The crystal structure of F_{28} TPP (Fig. 1) was determined, amongst others, in an attempt to correlate this porphyrin structure with the unusual spectroscopic data observed.



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